CLAIMS

What is claimed is:

- 1. An isolated polypeptide molecule comprising a contiguous sequence of 14 amino acids of SEQ ID NO:2.
- 2. An isolated polypeptide molecule according to claim 1, wherein the polypeptide molecule comprises residues 437 to 450 of SEQ ID NO:2.
- 3. An isolated polypeptide molecule according to claim 1 wherein the polypeptide molecule is between 82 and 232 amino acids in length.
- 4. An isolated polypeptide molecule according to claim 3 wherein the polypeptide molecule is residues 164 to 382 of SEQ ID NO:2.
- $\,$ 5. An isolated polypeptide molecule according to claim 3 wherein the polypeptide molecule is residues 383 to 464 of SEQ ID NO:2.
- 6. An isolated polypeptide molecule according to claim 3 wherein the polypeptide molecule is residues 465 to 696 of SEO ID NO:2.
- 7. A isolated polypeptide molecule selected from the group consisting of:
- a) a polypeptide molecule comprising residues 164 to 382 of SEO ID NO:2;
- b) a polypeptide molecule comprising residues 383 to 464 of SEO ID NO:2;

- c) a polypeptide molecule comprising residues 465 to 696 of SEO ID NO:2;
- d) a polypeptide molecule comprising residues 438 to 449 of SEO ID NO:2;
- e) a polypeptide molecule comprising residues 164 to 464 of SEQ ID NO:2;
- f) a polypeptide molecule comprising residues 164 to 696 of SEQ ID NO:2;
- g) a polypeptide molecule comprising residues 383 to 696 of SEQ ID NO:2;
- h) a polypeptide molecule comprising residues 164 to 449 of SEQ ID NO:2;
- i) a polypeptide molecule comprising residues 438 to 696 of SEQ ID NO:2; and
- j) a polypeptide molecule comprising residues 1 to 696 of SEQ ID NO:2.
- 8. An isolated polynucleotide molecule encoding a polypeptide molecule, wherein the polypeptide molecule comprises a contiguous sequence of 14 amino acids of SEQ ID NO:2.
- 9. An isolated polynucleotide molecule according to claim 8, wherein the polypeptide molecule comprises residues 437 to 450 of SEQ ID NO:2.
- 10. An isolated nucleotide molecule according to claim 8, wherein the polypeptide molecule is between 82 and 232 amino acids in length.
- 11. An isolated polynucleotide molecule according to claim 10, wherein the polypeptide molecule is residues 164 to 382 of SEQ ID NO:2.

- 12. An isolated polynucleotide molecule according to claim 10, wherein the polypeptide molecule is residues 383 to 464 of SEO ID NO:2.
- 13. An isolated polynucleotide molecule according to claim 10, wherein the polypeptide molecule is residues 465 to 696 of SEQ ID NO:2.
- 14. A isolated polynucleotide molecule encoding a polypeptide molecule, wherein the polypeptide molecule is selected from the group consisting of:
- a) a polypeptide molecule comprising residues 164
 to 382 of SEQ ID No:2;
- b) a polypeptide molecule comprising residues 383 to 464 of SEO ID NO:2;
- c) a polypeptide molecule comprising residues 465 to 696 of SEQ ID NO:2;
- d) a polypeptide molecule comprising residues 438 to 449 of SEO ID NO:2;
- e) a polypeptide molecule comprising residues 164 to 464 of SEQ ID NO:2;
- f) a polypeptide molecule comprising residues 164 to 696 of SEQ ID NO:2;
- g) a polypeptide molecule comprising residues 383 to 696 of SEQ ID NO:2;
- h) a polypeptide molecule comprising residues 164 to 449 of SEQ ID NO:2;
- i) a polypeptide molecule comprising residues 438 to 696 of SEQ ID NO:2; and
- j) a polypeptide molecule comprising residues 1 to 696 of SEO ID NO:2.
- 15. An isolated polynucleotide encoding a fusion protein having a first segment and a second segment, wherein

the first segment comprises a first polypeptide encoding a polypeptide having a protease domain and the second segment comprises a second polynucleotide encoding a polypeptide that has a contiguous sequence of 14 amino acids between residues 383 and 464 of SEQ ID NO:2, and wherein the first segment is positioned amino-terminally to the second segment.

- 16. An isolated polynucleotide according to claim 15, wherein the protease domain is selected from the group consisting of;
- a) a protease domain that is a member of the Disintegrin Proteases; and
- b) a protease domain that is at least 80% identical to amino acid residues 164 to 382 of SEQ ID NO:2.
- 17. An isolated polynucleotide molecule encoding a polypeptide molecule wherein the polynucleotide molecule is selected from the group consisting of:
- a) a polynucleotide molecule that encodes a polypeptide molecule that is at least 80 % identical to residues 383 to $\cdot 464$ of SEQ ID NO:2; and
- $\mbox{b)} \qquad \mbox{a polynucleotide molecule that is complementary} \\ \mbox{to a).}$
- 18. An isolated polynucleotide molecule according to claim 17 wherein the polynucleotide molecule is selected from the group consisting of:
- a) a polynucleotide molecule that encodes a polypeptide molecule that is at least 80 % identical to residues 383 to 696 of SEQ ID NO:2; and
- $\mbox{b)} \quad \mbox{a polynucleotide molecule that is complementary} \\ \mbox{to a).}$

- 19. An isolated polynucleotide molecule according to claim 17, wherein the polynucleotide molecule is selected from the group consisting of:
- a) a polynucleotide molecule that encodes a polypeptide molecule that is at least 80 % identical to residues 1 to 696 of SEQ ID NO:2; and
- $\mbox{b)} \qquad \mbox{a polynucleotide molecule that is complementary} \\ \mbox{to a).}$
- 20. An expression vector comprising the following operably linked elements:
 - a) a transcription promoter;
- b) a DNA segment encoding the polypeptide of claim 1; and
 - c) a transcription terminator.
- 21. An expression vector of claim 20 wherein the DNA segment further encodes an affinity tag.
- 22. A cultured cell into which has been introduced an expression vector according to claim 21, wherein said cell expresses the polypeptide encoded by the DNA segment.
- 23. A method of producing a polypeptide comprising culturing a cell according to claim 22, whereby said cell expresses the polypeptide encoded by the DNA segment; and recovering the polypeptide.
- 24. A method for modulating cell-cell interactions by combining the polypeptide according to claim 1, with cells in vivo and in vitro.

- 25. A method for modulating cell-cell interactions according to claim 24, whereby the cells are derived from tissues selected from the group consisting of:
 - a) tissues from heart;
 - b) tissues from brain;
 - c) tissues from spinal cord; and
 - d) tissues from skeletal muscle.
- 26. An isolated polypeptide molecule comprising a contiguous sequence of amino acids, wherein the contiguous sequence of amino acids is selected from the group consisting of:
 - a) SEQ ID NO:7;
 - b) SEQ ID NO:8;
 - c) SEQ ID NO:9;
 - d) SEQ ID NO:10; and
 - e) SEQ ID NO:11.
- 27. An isolated polynucleotide molecule encoding the isolated polypeptide molecule of claim 26.